

Amendments to the Specification:

Replace the paragraph beginning at page 1, line 4, with the following amended paragraph:

This application is a divisional of U.S.S.N. 09/855,266 filed May 14, 2001, now U.S. Patent No. 6,784,284, which is a divisional of U.S. Patent Application No. 09/411,722, filed October 1, 1999, now U.S. Patent No. 6,271,366, which is a continuation-in-part of PCT/JP98/01511 filed April 1, 1998, and claims priority from Japanese Application No. 9/99653 filed April 1, 1997.

Replace the paragraph beginning at page 21, line 15, with the following amended paragraph:

As used herein, "percent identity" of two amino acid sequences or of two nucleic acids is determined using the algorithm of Karlin and Altschul (Proc. Natl. Acad. Sci. USA 87:2264-2268, 1990), modified as in Karlin and Altschul (Proc. Natl. Acad. Sci. USA 90:5873-5877, 1993). Such an algorithm is incorporated into the NBLAST and XBLAST programs of Altschul et al. (J. Mol. Biol. 215:403-410, 1990). BLAST nucleotide searches are performed with the NBLAST program, score=100, wordlength=12 to obtain nucleotide sequences homologous to a nucleic acid molecules of the invention. BLAST protein searches are performed with the XBLAST program, score=50, wordlength=3 to obtain amino acid sequences homologous to a reference polypeptide (e.g., SEQ ID NO:2). To obtain gapped alignments for comparison purposes, Gapped BLAST is utilized as described in Altschul et al. (Nucleic Acids Res. 25:3389-3402, 1997). When utilizing BLAST and Gapped BLAST programs, the default parameters of the respective programs (e.g., XBLAST and NBLAST) are used. See <http://www.ncbi.nlm.nih.gov>.

Replace the title with the following amended title:

ISOLATED ANTIBODIES TO A SECRETORY MEMBRANE PROTEIN

Applicant : Naoki Kimura et al.
Serial No. : 10/802,332
Filed : March 16, 2004
Page : 3 of 15

Attorney's Docket No.: 14875-040003 / C1-806PCT-USD2

Replace the sequence listing with the new sequence listing attached.